

S6: PCA map of the protein expression data. Included are 69 identified proteins that were able to group normal endometrium (black), diploid- (red) and aneuploid- (green) endometrioid-carcinoma and aneuploid UPSC (blue) after outlier removal. X- and y-axis show first and second principal components that explain 68.2% of the variance in the data.

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